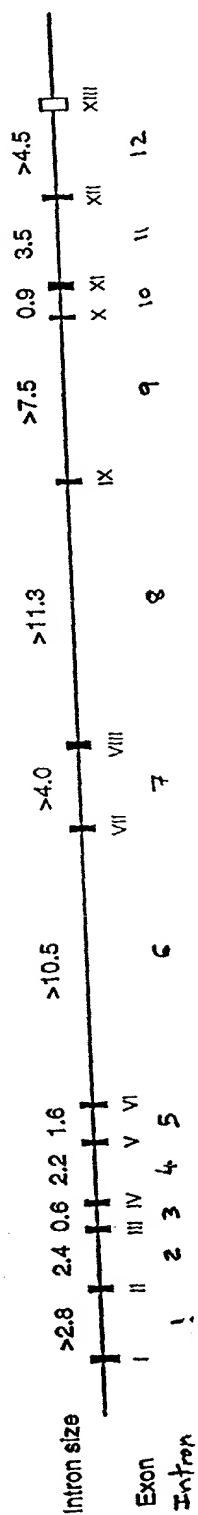


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Figure 1



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Figure 2A

promoter and exon 1

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GGTGAGGGGCTCAGAGGGGAGAGCTGGGAGGAGGGGAGA  
CATAGGTGGGGGAAGGGGTAGGAGAAAAGGGGAAGGGAGC  
AAGAGGGTGAGGGGCACCAGGCCCCATAGACGTTTGGC  
TCAGCGGCCACGAGGCTTCATCAGCTCCCGCCCCAAAAC  
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TCAGCGCCCCGCCCCGTCCCCGCCCCGACCCCGCCCCGG  
GCCCGCTCAGGCCCCGCCCCGTGCCCGCCGAATCCTGAAG  
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CCTGCGTGCCCGGAGTCCCCGCCTGTGTCTCTCTGTCTG  
CCGTCCCCGTCTCCTGCCAGGCGCGGAGCCCTGCCGAGCC  
GCGGGTGGGCCCCAGGCGCGCAGACATGCTGCTGCTCCGC  
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GCTACTGTGCGCTGTGCTGGGCGCTGTCATGATCGTGAT  
GGTGCCGTGCTCATCAAGCAGCAGGTCCTTAAG

A

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GCGGTGGGTGGGCGACCGCAGCGGAATCGGCGCCCGGGC  
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AGGGGCTGCTGCCCGCCTCCCCACCACCCTCACC

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[illegible]

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GGCAGCTGTTCCGGAGCCTTGTGGTGGGGCGTGGGGCC  
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CTGCAG

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CGCTGGGCATGGGACGGGTCTCANAGTGGACGGGATG  
GGGAGGCTGCTGACTGACCCCCAAACATTGTTCCGGAA  
GCACGCAACTCATAGTCGGGGTAAGTGCTACTCCCAAAA  
AAGTTTGCCT

CATGTCCCTGCAGTGGGCAGGCAGCGGGAGGGACAGACTT  
GGCGAAGGGGCCGAGCTCAGCTTTGGCTGTGGGGCCGGA  
GGTGTGCACAGACGTCCAGGGCCCCTGGTTCCCAGGCAG  
GCATTGCAGGCGAGTAGAAGGGAAACGTCCCATGCGAG  
CGGGGCGGGGCGTCTGACCCACTGGCTTCCCCCACAG

GTGAGGCTGCCCTGTGGCCACGCCGCTCGCACCCCTGA  
CCTCGTCCCTGTCTCTCCTCCCGCTGCCCTTGTG  
CAGAGAGCAGTCCCTGAGGTGGTCGGAGCGTGGGGACTC  
ACGCCCTGTGGGTGGCTTTCGGCCCTGTGCTGTCTCCAC  
CACCCCA

## Figure 2C

### exon 4

GGTGGTTCTGGTGTCCCAGATGCCCCACGTGGCCACTCC  
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CTTTGCTGTGAGGCCAGCTGGGGCCAAGGGAGGATG  
GGCCAGCCACGTCCAGCCTCTGACACTAGTGTCCCTTCG  
CCTTGCAG

GGTGCGGCGGTGATGATGGAGAATAAGCCCATGACCCTG  
AAGCTCATCATGACCTTGGCATTACACCACCTCGGCGAA  
CGTGCCCTTCATGAACCGCACTGTGGGTGAGATCATGTGG  
GGCTACAAGGACCCCTTGTGAATCTCATCAACAAGTACT  
TTCCAGGCATGTTCCTTCAAGGACAAGTTCGGATTAT  
TTGCTGAG

GTACGTGTGGCCTGGTGAGAAGCCAAAGATTCAGGCCTG  
TGTCTGTCTTCCCCTCACACAGCCTGGACACTGGTC  
ACCAGCTTGCTTTGTAGCTGGCTGGGGATCTAGTGGCTG  
TGGGTTGTAAGTGACTGAGAACCTGACTCAAACCGGCTT  
GAGTGAAA

### exon 5

CCTCTCGGTCCCCAGACACTGGGCATTTGGCAGTGAACC  
AGATGCTGGGGGCCCTGTCCTTCTGGTGGAGGGGGAGGA  
GGGCTCAGCCCAGAATGTTTCAGACCAGGCCGGCTCAA  
TGGCAGGCCTAAGCCTTACGATGCTGTTCCCTGCTGTGT  
CTGTAG

CTCAACAACCTCCGACTCTGGGCTCTTCACGGTGTTCACG  
GGGGTCCAGAACATCAGCAGGATCCACCTCGTGGACAAG  
TGGAACGGGCTGAGCAAG

GTGAGGGGCGAGAGGCGAGGGCCCCCTGTGCCAGGGAGA  
GGGGAGGGTGGGCCGCGCCATGGCTGCTCGGGAGTGGCA  
GGGACCAGAGAGCTCCTTCTTCTTTGTGCTGAAGAG  
GGTGCTGGGAGGATGAACACTCTTGAAGTTGGAGGAGGG  
ATTTTA

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T

## Figure 2D

exon 6

TCTCTGTGTGTCTACATAGCCTGCCCTCTTCCCACCGTG  
CCAGTATTGGGAATTGAGTGGCCGTGCGTGCACCAGGGT  
GAGTTAGGTGTGCAGCACCTGAGAGGGCTTATTAAGG  
GGCCTTGGCCCTACTGAGGGGTCTAGTCTGGATGCTTCC  
CCCCAG

GTTGACTTCTGGCATTCCGATCAGTGCAACATGATCAAT  
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GAGTCCCTCGCTGGAGTCTACAGCCCGAGGCCTGCCG

GTAATCACTGGGACTCGGGGCCTCCTGGGTTCCTGGGT  
AGCTCATGGCCAAATTCTGTGGTGTGGCTGTGCACCT  
GGAAAGCATTTTGACTCATCGTGGATTTGACTCAGTAG  
CCCTTGGCACCAGCTTGAATTCTCTTTGGTACACCACC  
AAAAGC

exon 7

GGAGGTGCGTGCAGCTCCGCGGGTGAGAGATGGGGGCGG  
TTTGGACCCGGGAGGTGGTAGCGCCCGTGGGGAGAAGTG  
GCTGGATCTGGGCAGCCTTTGGCAGGGCCTGGCTCTGGC  
CGCCGGGTCTGGGTGTCCCCTCTCATCCTGTCTGTCC  
CCTGCAG

ATCCATGAAGCTAATGTACAAGGAGTCAGGGGTGTTTGA  
AGGCATCCCCACCTATCGCTTCGTGGCTCCCCAAAACCT  
GTTTGCCAAACGGGTCCATCTACCCACCCAACGAAGGCTT  
CTGCCCCGTGCCTGGAGTCTGGAATTCAGAACGTCAGCAC  
CTGCAGGTTCA

GTACGTGCCGTCCCCCTGTTCTGGGATNGCCGGAGGGTGT  
TAGGTNTNGGGCACCTNANGGTTTATCTGCCCAATGCTG  
TCTGCTTAATCTCTGGCCTCTGTACTCTTGATAACC  
CATTAAGCCAAAAATATGATGCCTCTGGGACGATATCTG

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## Figure 2E

exon 8

TGGGGCTTTTACAGAATGGAGGAAGGGATCCTCTCT  
GTCGGGTATTATGGTCATCGCCACGGGGGTGCCGTGCAG  
ACCACAGCTCTGTGCAGACTTCCGGAGTGGCAGGACGTG  
CCAATATACTGTCGTTGTATGATGTCCCCCTCCCTGCCCT  
TGTGTAG

GTGCCCCCTGTTTCTCTCCCATCCTCACTTCCTCAACG  
CTGACCCGGTTCTGGCAGAAAGCGGTGACTGGCCTGCACC  
CTAACCCAGGAGGCACACTCCTTGTTCCTGGACATCCACC  
CG

GTGAGCCCCCTGCCATCCTCTGTGGGGGGTGGGTGATTCC  
TGGTTGGAGCACACCTGGCTGCCTCCTCTCTCCCCAG  
GCAGAGAGCTGCTGTGGGCTGGGGTGGTGGGAAGCCTGG  
CTTCTAGAATCTCGAGCCACCAAAGTTCCTTACT

exon 9

CCCCAGCCTGTGGCTTGTTTTAGGTAAGATACAAGCAAG  
CTCCACTGGGCAGTTAGCTGGGACGCCCACCCTCTTGAC  
TGGGACCAGGGAAAAGAAGGTGACTGTGTCCCTGGA  
GCTTGGGGGTGGCCAGTCTCCTCACTGTGTTTGTGCCG  
CAG

GTCACGGGAATCCCCATGAACTGCTCTGTGAAACTGCAG  
CTGAGCCTCTACATGAAATCTGTCCGAGGCATTGG

GTGAGTGGGGACTGGGAACCTGGGGCTGCATTGCTCATTG  
AGAGATTANGTGCTCAGTGCTCCAGTGTTCCCAGAC  
TCCCCTGACATACCCAGGAAACAGGGCATGGGGAAGGG  
AGAGGGTCCTATTGGGGGTGGAATCCAGTCCCTGCTGAT  
CTTCTC

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## Figure 2F

exon 10

ATGGCTCCTAAAGTGTTTCAGCTCATTGTTTATATTGG  
TGGTGAGGGTTTAGTGTGTGCAAAATTATACTAAACC  
TGTTTAGATGTTGTATTCAAGCAGAATTAGATCAAGTTT  
GGGTGTAAGACTTTGTTCCAACACCTATGTCTTGCTTAT  
TTCCAG

ACAAACTGGGAAGATTGAGCCTGTGGTCCTGCCGCTGCT  
CTGGTTTGCAGAG

GTAAGGGTGCGTTGGGCACAGCGTCGGGGGCTTTTGTTA  
ATAGCCAATGTGGGCATTTGAGGCAGGAGGCGGGGGG  
AGCACCTTGTAAGAAAGGGAGAGGGCTGAGCCAGGGTAAC  
CGGACTGTTACATGGACCAGCGTATCATACACTTCACCC  
TGTC

exon 11

CCTGGAGGGAGGAGGTCCCTGGCAGGCTCCAACACATGC  
TTTAGCCGGGAAGCTTGAGGTGGGGAAAAGCTGAGGCGG  
GCACAGAGGAAGGTGTGGGTGGCATCTGCGCTGTAG  
CCCGCAGCGTGGCGCCCCAGCTCATGTGTTGTGTCATTCT  
GTCTCCTCAG

AGCGGGGCCATGGAGGGGGAGACTCTTCACACATTCTAC  
ACTCAGCTGGTGTTGATGCCCAAGGTGATGCACTATGCC  
CAGTACGTCCCTCCTGGCGCTGGGCTGCGTCCTGCTGCTG  
GTCCCTGTCATCTGCCAAATCCGGAGCCAA

GTAGGTGCTGGCCAGAGGGCAGCCCGGGCTGACAGCCAT  
TCGCTTGCCTGCTGGGGGAAAGGGGCCTCAGATCGGACC  
CTCTGGCCAACCGCAGCCTGGAGCCCACCTCCAGCAG  
CAGTCCTGCGTCTCTGCCGGAGTGGGAGCGGTCACTGCT  
GGGGG

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## Figure 2G

exon 12

CCCCACATCTCAGCCACCTGCAATCGTTGAGGGTTGTTG  
GACTCTAAACTTATGTGCCTTTCCTGTTTCCTCTTTGCC  
TTTTGCAAATTGAAGAACCGTGTA AAACCATTTTTAT  
GTGGCTTCAACGTCAACTATAAATTAGCTTGGTTATCTT  
CTAG

GAGAAATGCTATTTATTTGGAGTAGTAGTAAAAAGGGC  
TCAAAGGATAAGGAGGCCATTCAGGCCTATTCTGAATCC  
CTGATGACATCAGCTCCCAAGGGCTCTGTGCTGCAGGAA  
GCAAAACTGTAG

GTGGGTACCAGGTAATGCCGTGCGCCTCCCCGCCCCCTC  
CCATATCAAGTAGAATGCTGGCGGCTTAAACATTTGGG  
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TTGGAGAAA

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Figure 3A

ACCGTGCTCTGGGGCTGGGTGCCCGGAGTCCCGGCTGTGTCTCTGTGCGCGTCCCGCTCTCTGCCAGGCGGG 79

GAGCCCTGCGAGCGCGGGTGGGCCCCAGGCGGCGAGAC ATG GGC TGC TCC GCC AAA GCG GCG TGG GCT 10 148

A G A L G V A G L L C A V L G A V M I V 30  
GCC GGG GCG CTG GGC GTC GCG GGG CTA CTG TGC GCT GTG CTG GGC GCT GTC ATG ATC GTG 208

exon 1 → exon 2

H V P S L I K Q Q V L K N V R I D P S S 50  
ATG GTG CCG TCG CTC ATC AAG CAG CAG GTC CTT AAG AAC GTG GCG ATC GAC CCC AGT AGC 268

L S F N M W K E I P I P F Y L S V Y F F 70  
CTG TCC TTC AAC ATG TGG AAG GAG ATC CCT ATC CCC TTC TAT CTC TCC GTC TAC TTC TTT 328

D V M N P S E I L K G E K P Q V R E R G 90  
GAC GTC ATG AAC CCC AGC GAG ATC CTG AAG GGC GAG AAG CCG CAG GTG GCG GAG GCG GGG 388

exon 3

P Y V Y R E F R H K S N I T F N N N D T 110  
CCC TAC GTG TAC AGG GAG TTC AGG CAC AAA AGC AAC ATC ACC TTC AAC AAC AAC GAC ACC 448

V S F L E Y R T F Q F Q P S K S H G S E 130  
GTG TCC TTC CTC GAG TAC CCG ACC TTC CAG TTC CAG CCC TCC AAG TCC CAC GGC TCG GAG 508

exon 4

S D Y I V M P N I L V L G A A V H M E N 150  
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K P M T L K L I H T L A F T T L G E R A 170  
AAG CCC ATG ACC CTG AAG CTC ATC ATG ACC TTG GCA TTC ACC ACC CTC GGC GAA CGT GCC 628

F M N R T V G E I M W G Y K D P L V N L 190  
TTC ATG AAC CGC ACT GTG GGT GAG ATC ATG TGG GGC TAC AAG GAC CCC CTT GTG AAT CTC 688

exon 5

I N K Y F P G M F P F K D K F G L F A E 210  
ATC AAC AAG TAC TTT CCA GGC ATG TTC CCC TTC AAG GAC AAG TTC GGA TTA TTT GCT GAG 748

L N N S D S G L F T V F T G V Q N I S R 230  
CTC AAC AAC TCC GAC TCT GGG CTC TTC ACG GTG TTC ACG GGG GTC CAG AAC ATC AGC AGG 808

exon 6

I H L V D K W N G L S K V D F W H S D Q 250  
ATC CAC CTC GTG GAC AAG TGG AAC GGG CTG AGC AAG GTT GAC TTC TGG CAT TCC GAT CAG 868

C N M I N G T S G Q M W P P F M T P E S 270  
TGC AAC ATG ATC AAT GGA ACT TCT GGG CAA ATG TGG CCG CCC TTC ATG ACT CCT GAG TCC 928

exon 7

S L E F Y S P E A C R S M K L M Y K E S 290  
TCG CTG GAG TTC TAC AGC CCG GAG GCC TGC CGA TCC ATG AAG CTA ATG TAC AAG GAG TCA 988

G V F E G I P T Y R F V A P K T L F A N 310  
GGG GTG TTT GAA GGC ATC CCC ACC TAT CCG TTC GTG GCT CCC AAA ACC CTG TTT GCC AAC 1048

G S I Y P P N E G F C P C L E S G I Q N 330  
GGG TCC ATC TAC CCA CCC AAC GAA GGC TTC TGC CCG TGC CTG GAG TCT GGA ATT CAG AAC 1108

exon 8

V S T C R F S A P L F L S H P H F L N A 350  
GTC AGC ACC TGC AGG TTC AGT GCC CCC TTG TTT CTC TCC CAT CCT CAC TTC CTC AAC GC 1168

D P V L A E A V T G L H P N Q E A H S L 370  
GAC CCG GTT CTG GCA GAA GCG GTG ACT GCC CTG CAC CCT AAC CAG GAG GCA CAC TCC TTG 1228

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Figure 3B

F L D I H P V T G I P M N C S V K L Q L 390  
 TTC CTG GAC ATC CAC CCG GTC ACG GGA ATC CCC ATG AAC TGC TCT GTG AAA CTG CAG CTG 1288  
 S L Y M K S V A G I G Q T G K I E P V V 410  
 AGC CTC TAC ATG AAA TCT GTC GCA GGC ATT GGA CAA ACT GGG AAG ATT GAG CCT GTG GTC 1348  
 L P L L W F A E S G A M E G E T L H T F 430  
 CTG CCG CTG CTC TGG TTT GCA GAG AGC GGG GCC ATG GAG GGG GAG ACT CTT CAC ACA TTC 1408  
 Y T Q L V L M P K V M H Y A Q Y V L L A 450  
 TAC ACT CAG CTG GTG TTG ATG CCC AAG GTG ATG CAC TAT GCC CAG TAC GTC CTC CTG GCG 1468  
 L G C V L L L V P V I C Q I R S Q E K C 470  
 CTG GGC TGC GTC CTG CTG CTG GTC CCT GTC ATC TGC CAA ATC CGG AGC CAA GAG AAA TGC 1528  
 Y L F W S S S K K G S K D K E A I Q A Y 490  
 TAT TTA TTT TGG AGT AGT AGT AAA AAG GGC TCA AAG GAT AAG GAG GCC ATT CAG GCC TAT 1588  
 S E S L M T S A P K G S V L Q E A K L \* 510  
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Figure 4

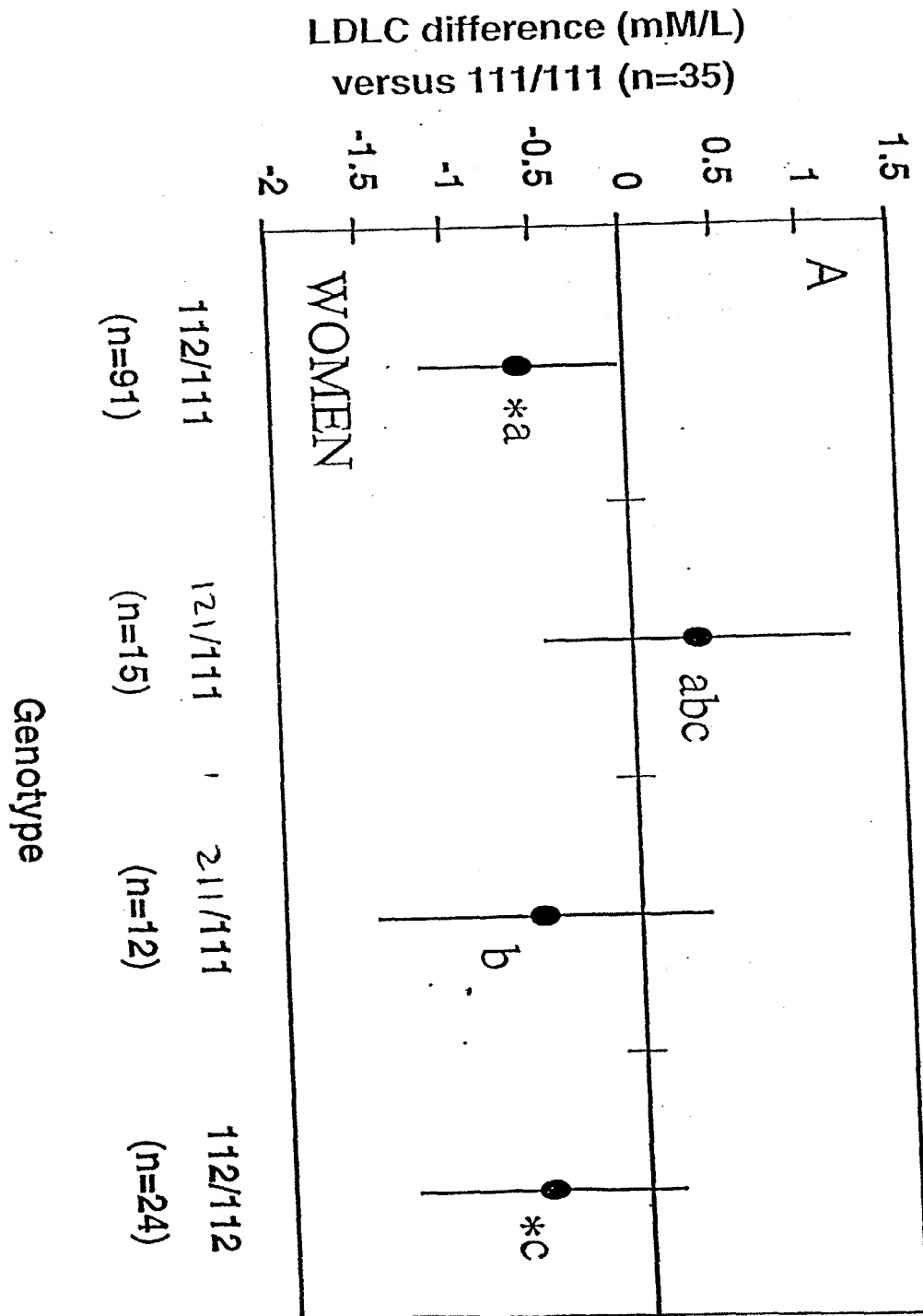
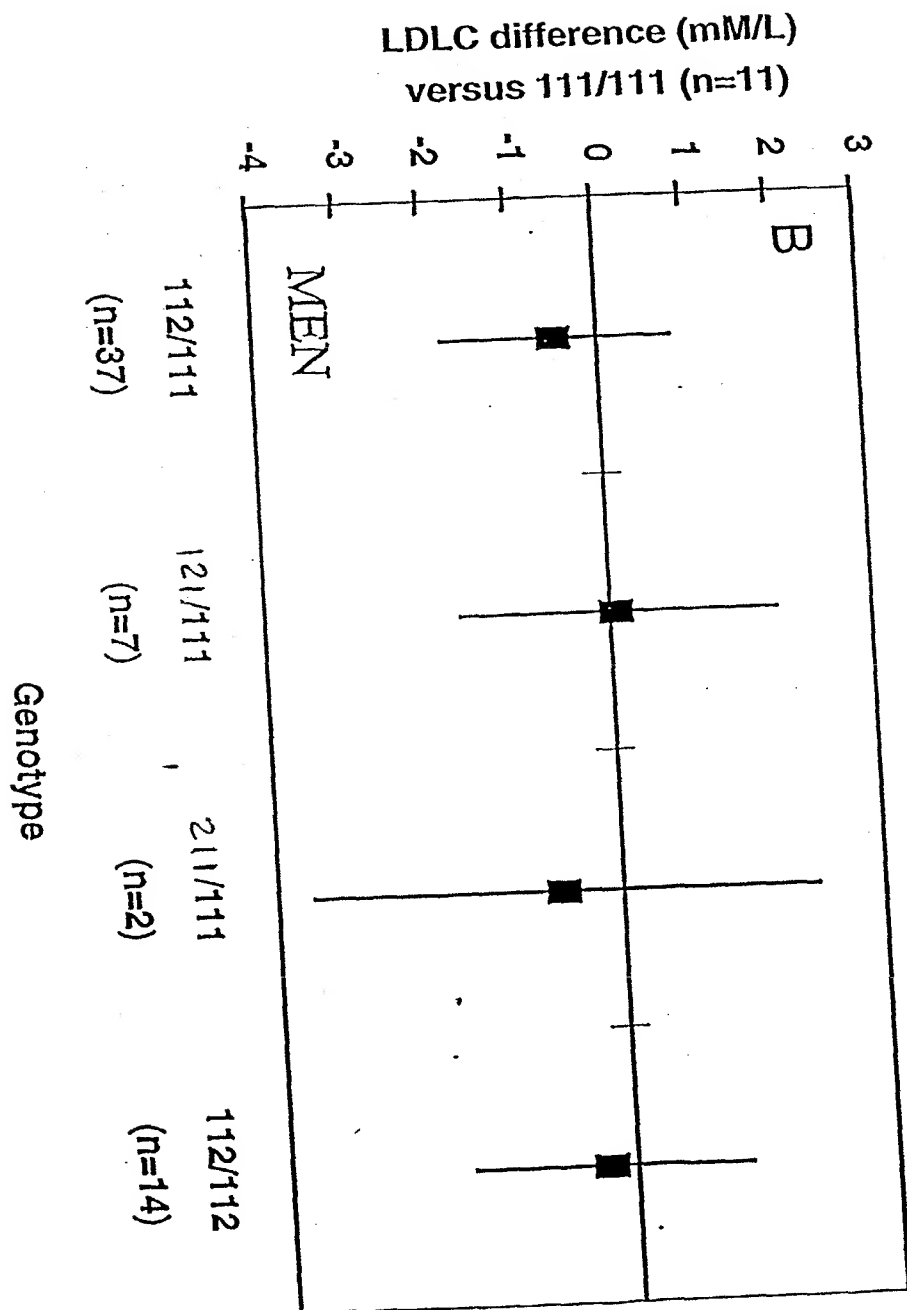
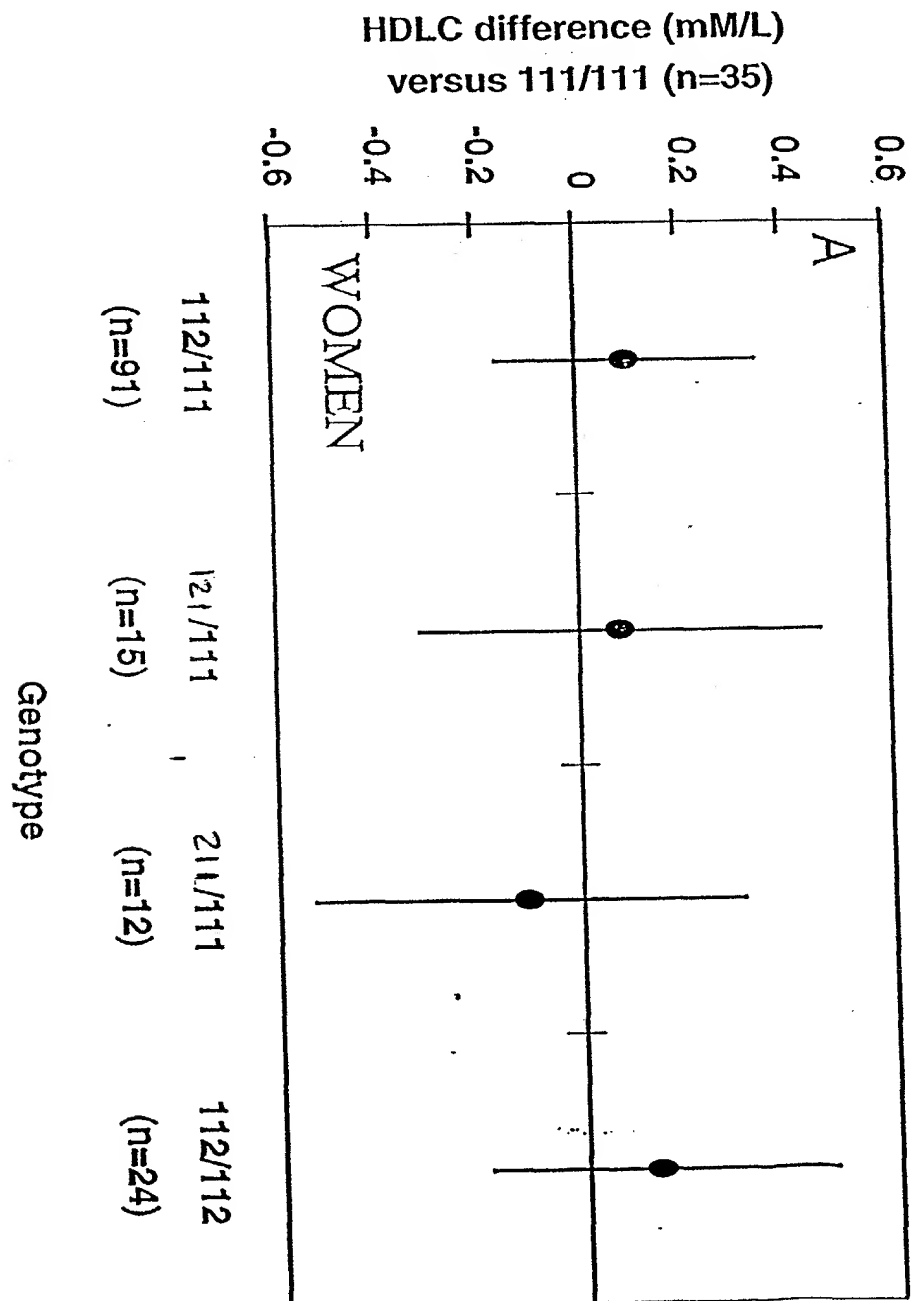


Figure 5



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Figure 6



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Figure 7

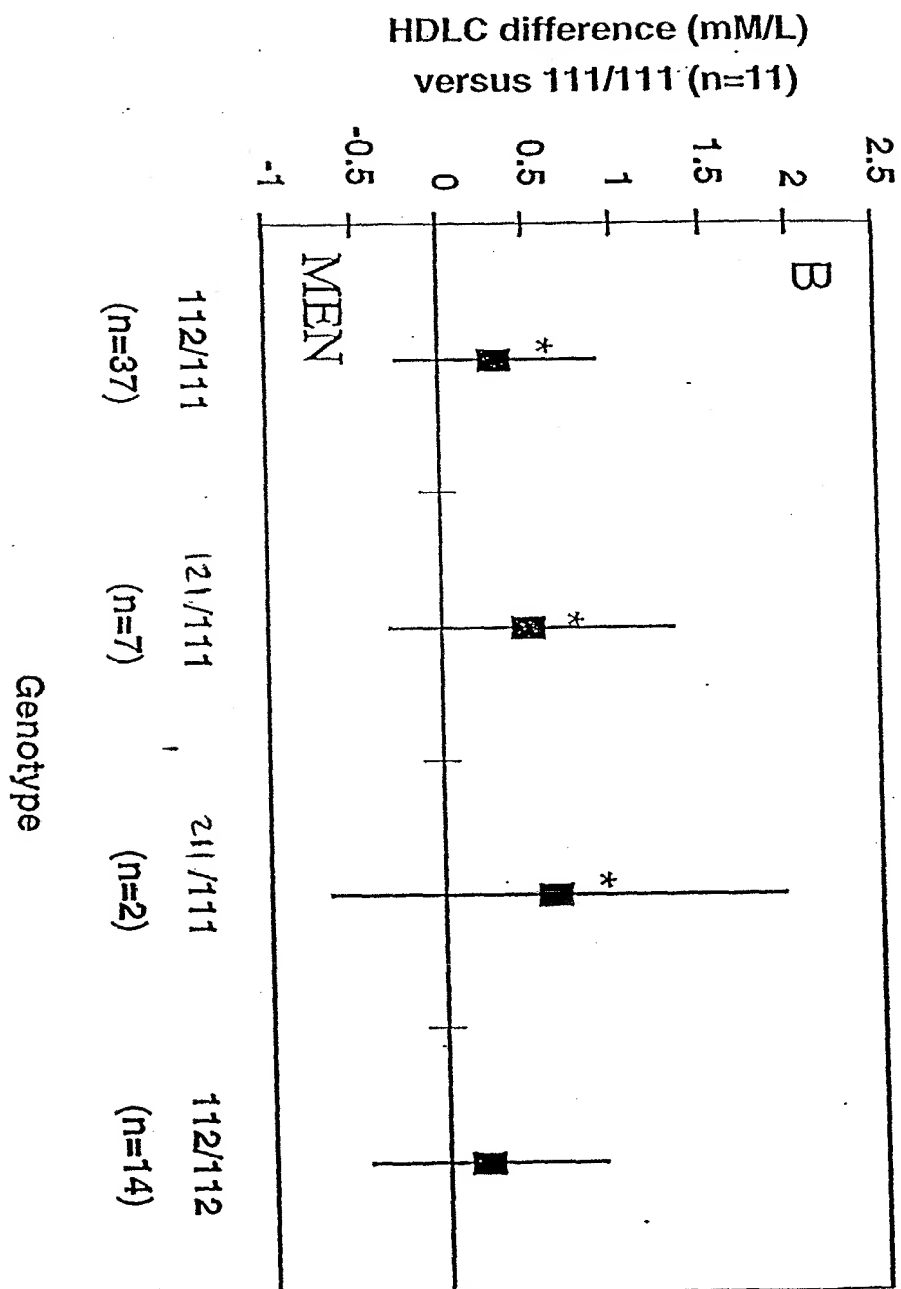
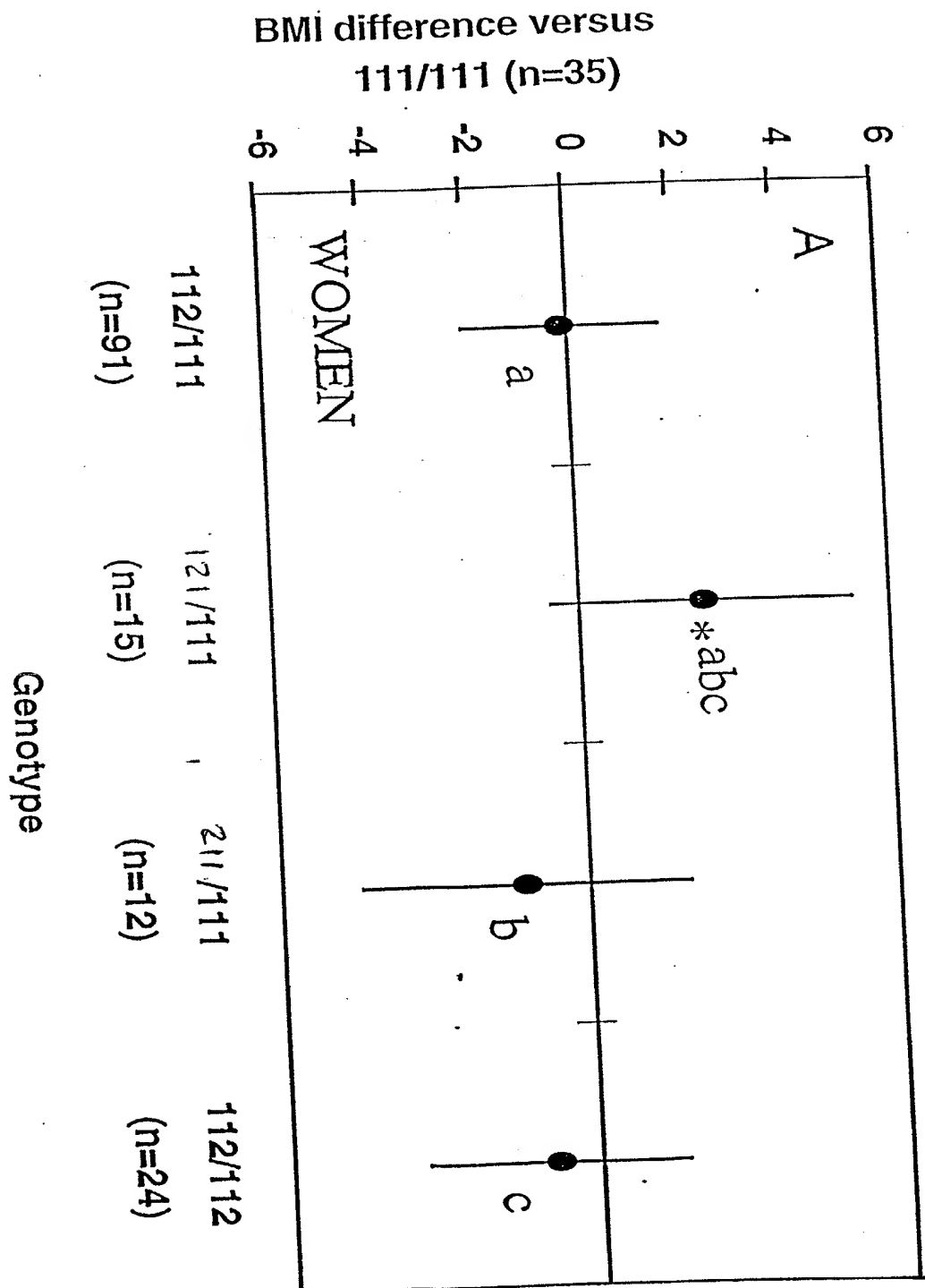
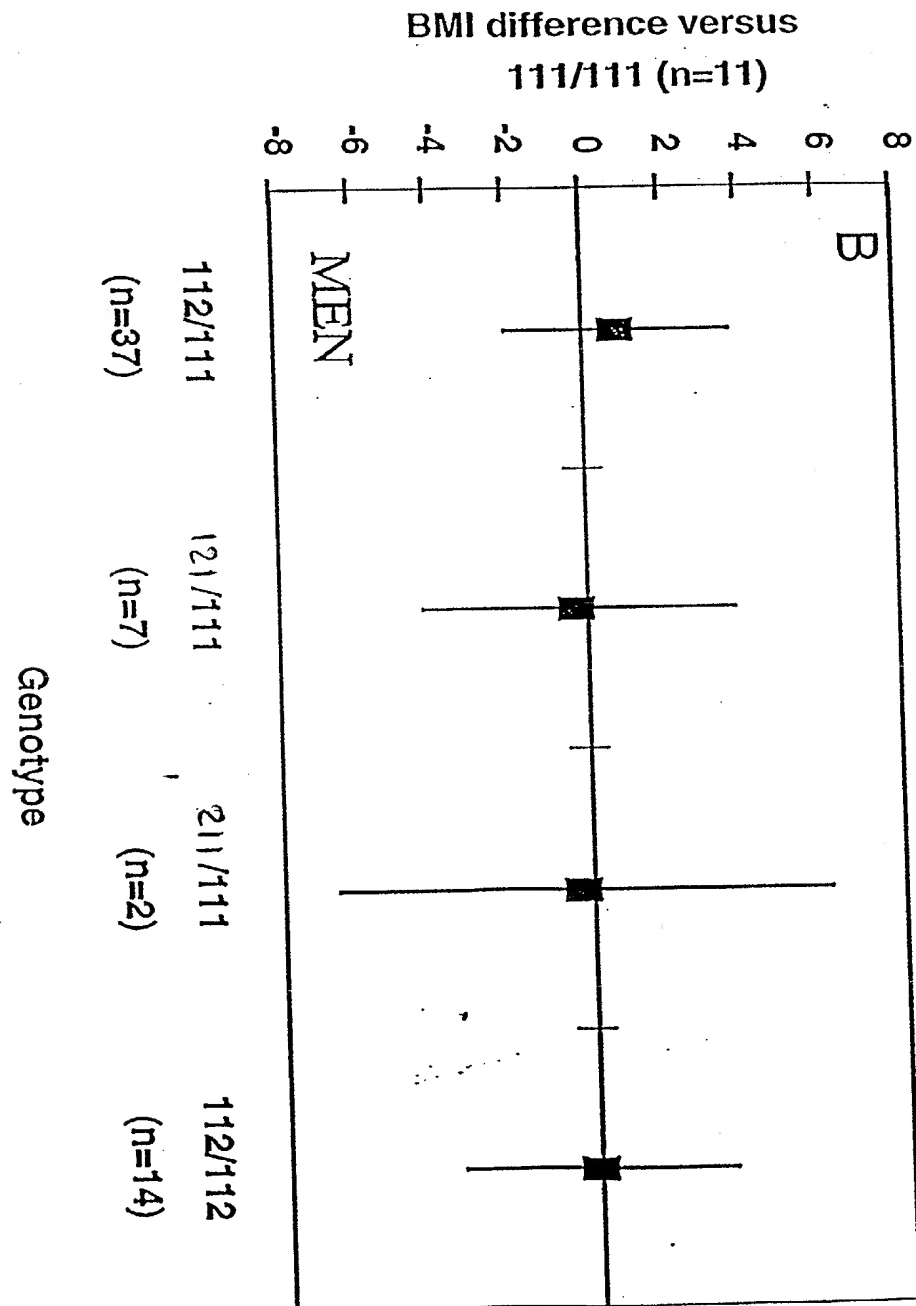


Figure 8



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Figure 9



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